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Early accumulation of intracellular fibrillar oligomers and late congophilic amyloid angiopathy in mice expressing the Osaka intra-A β APP mutation

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Pathogenic amyloid- β peptide precursor (APP) mutations clustered around position 693 of APP—position 22 of the A β sequence—are commonly associated with congophilic amyloid angiopathy (CAA) and intracerebral hemorrhages. In contrast, the Osaka (E693 Δ) intra-A β APP mutation shows a recessive pattern of inheritance that leads to AD-like dementia despite low brain amyloid on *in vivo* positron emission tomography imaging. Here, we investigated the effects of the Osaka APP mutation on A β accumulation and deposition *in vivo* using a newly generated APP transgenic mouse model (E22 Δ A β) expressing the Osaka mutation together with the Swedish (K670N/M671L) double mutation. E22 Δ A β mice exhibited reduced α -processing of APP and early accumulation of intraneuronal fibrillar A β oligomers associated with cognitive deficits. In line with our *in vitro* findings that recombinant E22 Δ -mutated A β peptides form amyloid fibrils, aged E22 Δ A β mice showed extracellular CAA deposits in leptomeningeal cerebellar and cortical vessels. *In vitro* results from thioflavin T aggregation assays with recombinant A β peptides revealed a yet unknown anti-amyloidogenic property of the E693 Δ mutation in the heterozygous state and an inhibitory effect of E22 Δ A β 42 on E22 Δ A β 40 fibrillogenesis. Moreover, E22 Δ A β 42 showed a unique aggregation kinetics lacking exponential fibril growth and poor seeding effects on wild-type A β aggregation. These results provide a possible explanation for the recessive trait of inheritance of the Osaka APP mutation and the apparent lack of amyloid deposition in E693 Δ mutation carriers.

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Introduction

Extracellular deposition of fibrillar amyloid- β (A β) peptide as amyloid plaques and congophilic amyloid angiopathy (CAA) is considered a cardinal neuropathological feature of Alzheimer's disease (AD).^{1,2} According to the amyloid cascade hypothesis, soluble and fibrillar A β species have a central role in the pathogenesis of AD and start to accumulate within the brain years before cognitive decline and dementia symptoms are observed.³ Evidence supporting the amyloid cascade hypothesis comes from several sources including genetic studies of familial AD cases carrying mutations in the amyloid- β peptide precursor (APP) gene and the *presenilin* genes.^{3,4} Pathogenic mutations in the APP gene have been shown to influence the metabolism of the A β peptide in various ways. The Swedish double mutation (K670N/M671L), located upstream of the A β N terminus adjacent to the β -cleavage site, results in an increased production of both A β 40 and A β 42 species,^{5–7} whereas mutations located at the γ -cleavage site of APP cause an increase of the A β 42/A β 40 ratio and, as a consequence of this, result in increased A β aggregation and deposition (for a review see Goate⁸). The recently discovered

E693 Δ Osaka mutation in a Japanese pedigree⁹ is one of the six so-called intra-A β mutations clustered around the hydrophobic core of the A β sequence. Position 693 seems to be a critical site involved in pathogenic aggregate formation since mutations at or (\pm 1) around this site, including the Dutch (E693Q),¹⁰ Flemish (E692G),¹¹ Italian (E693K),¹² Iowa (D694N)¹³ and Arctic (E693G)¹⁴ mutations, have been reported to result in an increase in total A β production¹⁵ and—with the exception of the Flemish mutation—enhance A β aggregation and toxicity.^{16,17} Interestingly, all currently known intra-A β APP mutations—with the exception of E693 Δ —have previously been shown to be vasculotropic and are characterized neuropathologically by prominent vascular amyloid deposition.¹⁸

Although neuropathological data have not been reported to date, homozygous carriers of the recessive Osaka APP mutation are believed to develop an AD-like clinical phenotype in the absence of relevant extracellular amyloid deposition as revealed by a very low signal on amyloid positron emission tomography imaging.⁹ *In vitro* experiments demonstrated enhanced oligomerization but no fibrillization of synthetic

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E22 Δ A β 40 and E22 Δ A β 42 preparations, suggesting that AD-like symptoms may be caused by the presence of synaptotoxic A β oligomers, rather than fibrillar A β , in the affected patients.⁹ Consistent with these findings, synthetic E22 Δ A β 42 potently inhibited hippocampal long-term potentiation^{9,19} and induced synapse loss in mouse hippocampal slices.²⁰ Further cell culture experiments and results from the recently reported E693 Δ transgenic mouse model revealed enhanced accumulation of intraneuronal A β oligomers as a prominent feature of the Osaka APP mutation.^{21–23} E693 Δ transgenic mice start to accumulate intraneuronal A β aggregates at an age of 8 months and are completely devoid of extracellular amyloid deposits up to an age of 24 months.²² The apparent lack of extracellular amyloid deposition in these mice has been suggested to be in line with the initial *in vitro* findings with synthetic A β preparations that E22 Δ -mutated A β peptides do not form amyloid fibrils.⁹

However, follow-up studies with recombinant A β preparations revealed that both E22 Δ A β 40 and E22 Δ A β 42 readily formed amyloid fibrils *in vitro*.²⁴ Based on these findings, we hypothesized that E22 Δ -mutated A β may, in principle, also form amyloid fibrils *in vivo* and generated a novel APP transgenic mouse line (E22 Δ A β) to investigate the effects of the E693 Δ mutation on amyloid accumulation and deposition *in vivo*. In line with our recent *in vitro* findings,²⁴ aged E22 Δ A β mice were characterized by late extracellular amyloid deposition in the leptomeningeal vasculature at 24 months of age, which was preceded by an early accumulation of intracellular oligomeric A β species already at an age of 3 months. The results of this study provide strong evidence that E22 Δ -mutated A β species are fibrillogenic and can deposit extracellularly *in vivo* as CAA, thus placing the E693 Δ Osaka mutation on the list of other vasculotropic intra-A β APP mutations.

Materials and methods

Animals. The newly generated E22 Δ A β mice express the human APP695 isoform containing the Swedish (K670N + M671L) and Osaka (E693 Δ) mutations. The mutations were generated by site-directed mutagenesis of pGEM-9zf(–)-huAPP695. The cDNA was inserted into pMoPrP-Xho,²⁵ and the construct was sequenced. After removal of the vector sequence, the linear construct was injected into the pronuclei of fertilized zygotes of B6D2F1 mice. Founders were screened for transgene expression by tail polymerase chain reaction and western blot analysis, and the line used in this project was expanded on the hybrid background of C57Bl/6 and DBA/2 (B6D2). For all behavioral, biochemical and histological analyses, offspring from the B6D2 generation backcrossed once with pure C57Bl/6 was used. Mice were kept on a 12 h light/dark cycle at 22 °C. Food pellets and water were available *ad libitum*.

For histological and western blot analyses, mice at different ages were anesthetized and transcardially perfused with ice-cold phosphate-buffered saline. One hemibrain was dissected into cortex and hippocampus, frozen in liquid nitrogen and stored at –80 °C. The other hemibrain was postfixed in 4% (w/v^{–1}) paraformaldehyde in phosphate-buffered saline overnight at 4 °C and embedded in paraffin.

All animal experiments were performed in compliance with Swiss national guidelines and were approved by the veterinarian authorities of the Canton of Zurich.

Protein extracts and western blotting. Brain tissues were homogenized with a glass tefflon homogenizer in a sixfold wet weight amount of buffer A containing 100 mM Tris-HCl, 150 mM NaCl, Complete Protease Inhibitor Cocktail (Roche Diagnostics, Rotkreuz, Switzerland) and Phosphatase Inhibitor Cocktails 1 + 2 (Sigma-Aldrich, Buchs, Switzerland). After centrifugation at 100 000 *g* for 1 h, supernatants were collected (= Tris fraction) and pellets were rehomogenized in buffer A containing 1% Triton X-100. Centrifugation at 100 000 *g* was repeated and supernatants were again collected (= Triton fraction). The remaining pellets were rehomogenized in buffer A containing 2% sodium dodecyl sulfate (SDS). After an additional centrifugation step and collection of the supernatants (= SDS fraction), the resulting pellets were eventually dissolved in 70% formic acid (FA), sonicated for 30 s at 30% power, ultracentrifuged, supernatants extracted, lyophilized and reconstituted in buffer A containing 2% SDS for further analysis. Total protein concentrations were measured with the DC protein assay (Bio-Rad Labs, Gessier, Switzerland). Extracts were separated by SDS-polyacrylamide gel electrophoresis, blotted onto nitrocellulose, boiled for 5 min in phosphate-buffered saline and blocked in Tris-buffered saline containing 5% milk for 1 h at room temperature. Primary antibodies were incubated overnight at 4 °C (6E10 1:500; anti-C-terminal APP (Sigma, Buchs, Switzerland); 1:2000) and visualized by peroxidase-conjugated antibodies and ECL reactions (Amersham Biosciences, Otelfingen, Switzerland). Monoclonal anti- β -actin antibody (1:2000; Abcam, Cambridge, UK) was used as internal loading control and for normalization of densitometric analyses of the immunoreactive bands. Quantification of the immunoreactive bands was carried out by densitometry of the scanned films under conditions of non-saturated signal using the Image J software (rsb.info.nih.gov/ij/).

MSD analysis. A β fragments were measured in the above-mentioned brain homogenate fractions and plasma was measured using a MesoScale Discovery (MSD) 3plex multi-SPOT A β human kit (Gaithersburg, MD, USA) for A β ₃₈, A β ₄₀ and A β ₄₂, in accordance with the manufacturer's instructions. Human sAPP α levels were determined using an MSD 2plex kit (Gaithersburg, MD, USA), and human Swedish sAPP β levels were determined using an MSD 1plex kit (Gaithersburg, MD, USA), in accordance with the manufacturer's instructions. All reagents were provided with the kits containing a 96-well plate with two carbon electrodes precoated with analyte-specific capture antibodies. After 1 h of blocking with bovine serum albumin, plates were washed with Tris wash buffer and samples and standards added to the wells. Plates were sealed and incubated for 1 h on an orbital shaker (750 r.p.m.) at room temperature, followed by additional washing steps and incubation with the detection antibody for 1 h. For A β 3plex assay (Gaithersburg, MD, USA), standards and samples were added at the same time as the detection antibody, and plates were incubated for 2 h. At the end of the incubation period, plates were washed again and measured

on an MSD SECTOR Imager 6000 plate reader (Gaithersburg, MD, USA) after the addition of the MSD Read Buffer T (Gaithersburg, MD, USA). Raw data were measured as electrochemiluminescence (light) signal detected by photo-detectors. The MSD DISCOVERY WORKBENCH software (Version 3.0.17) (Gaithersburg, MD, USA) with Data Analysis Toolbox was used to calculate sample concentrations by comparing them against a standard curve.

Histological analysis. Histological stainings were carried out on 5 μ m paraffin brain sections by using standard published procedures. For the immunohistochemical detection of intraneuronal A β deposits, sections were boiled in 10 mM sodium citrate buffer (pH 6.0), followed by antigen retrieval with 95% FA for 5 min. The following antibodies were used for immunohistochemistry: 6E10 (1:500 dilution; Signet, Dedham, MA, USA) and β -amyloid antibody (1:200 dilution; Cell Signaling, Danvers, MA, USA) were used for the detection of pan-A β . Anti- β -amyloid protein (1–40) antibody (1:100; Sigma) and BA27 (Amyloid β -Protein Immunohisto-stain Kit; Wako, Wako Chemicals GmbH, Neuss, Germany) were used to specifically detect A β 40. β -Amyloid 42 Polyclonal Antibody (1:100; Signet) and BC05 (Amyloid β -Protein Immunohisto-stain Kit; Wako) were used to specifically detect A β 42. Polyclonal antibodies A11 and OC (provided by C Glabe, both 1:100) were used for the detection of prefibrillar and fibrillar A β oligomers, respectively. 11A1 (1:100; IBL Japan, Gunma, Japan) was used to detect A β with a conformational turn epitope between positions 22 and 23 of the A β sequence.²⁶ Anti-Amyloid Precursor Protein, C-Terminal antibody (1:200; Sigma) was used for the detection of full-length APP and APP C-terminal fragments. Secondary antibodies were obtained from Vector Laboratories, Burlingame, CA, USA (Vectastain ABC kits PK-6101 and PK-6102) for peroxidase diaminobenzidine stainings, and from Jackson Immunoresearch Laboratories (West Grove, PA, USA) for immunofluorescence.

Thioflavin S staining and Congo red stainings were performed according to standard protocols as described previously.²⁷

Recombinant A β production. Production of recombinant A β peptides (wild-type A β 40 and A β 42, E22 Δ A β 40 and A β 42, and E22G A β 40 and A β 42) was performed as described previously.²⁸ In brief, recombinant A β peptides were expressed under the control of the T7 promoter/lac operator in *Escherichia coli* BL21 (DE3) as fusion proteins to the peptide sequence (NANP)₁₉ with an N-terminal hexahistidine tag. Mutagenesis at codon 22 in both A β 1–40 and A β 1–42 was performed with the QuikChange site-directed mutagenesis kit (Stratagene, Basel, Switzerland). The correct genetic sequences of the constructs were verified by DNA sequencing. Cleavage of the fusion proteins (100 μ M) with 7.5 μ M tobacco etch virus protease was performed in 10 mM Tris-HCl (pH 8.0), 0.5 mM ethylenediaminetetraacetic acid and 1 mM dithiothreitol for 1 h at room temperature, followed by incubation at 4 °C overnight. The cleaved A β peptides precipitated during the cleavage reactions and were pelleted by centrifugation (4500 g, 20 min, 4 °C), dissolved in 6 M guanidinium chloride-HCl (pH 2.0) and purified via

reversed-phase high-performance liquid chromatography. Peptides were eluted with CAN, aliquoted in Protein LoBind Eppendorf tubes (Vaudaux-Eppendorf), lyophilized and stored at –80 °C. The high purity and identity of the peptides were verified by matrix-assisted laser desorption/ionization time-of-flight mass spectrometry using sinapinic acid as matrix.

Thioflavin T aggregation assays. Preparation of A β solutions and thioflavin T aggregation assays were performed as described earlier.^{24,28} In brief, A β variants were dissolved in 10 mM NaOH to concentrations of 100–150 μ M. A β concentrations were determined via the absorbance of A β at 280 nm in 10 mM NaOH (extinction coefficient at 280 nm and pH 12 corresponding to a single tyrosine residue: 1730 M^{–1} cm^{–1}). Stock solutions were kept on ice and were used for aggregation experiments within 6 h. Aggregation reactions were performed at 37 °C with 2.5 or 5 μ M A β (final concentration) in 10 mM H₃PO₄-NaOH (pH 7.4), 100 mM NaCl and 35 μ M thioflavin T in a volume of 1000 μ l in stirred quartz fluorescence cuvettes (1–0.4 cm). The thioflavin T concentration was determined via its extinction coefficient of 36 000 M^{–1} cm^{–1} at 412 nm. Aggregation reactions were started by a dilution of the A β stock solution in 10 mM NaOH (prepared and ultracentrifuged immediately before use) with an aggregation buffer mix, resulting in pH 7.4, and the final concentrations indicated above. Thioflavin T fluorescence emission at 482 nm (excitation at 440 nm; excitation and emission slit at 1.6 nm) was monitored on a Quantamaster (QM-7/2003) fluorescence spectrometer (Photon Technology International, Birmingham, NJ, USA).

Cognitive-behavioral testing. A battery of well-validated and carefully controlled tests was used to behaviorally assess mice for motoric and cognitive performance.²⁹ At the time of testing, mice were weighed and examined for general health measures to ensure that the mice were physically able to conduct the cognitive-behavioral test.²⁹

Locomotor activity in the open field test. According to published procedures, mice were placed in the center of a brightly-lit white Plexiglas box (50 \times 50 cm²), and their movements (distance traveled and average speed) were tracked using ANY-maze video tracking software (Stoelting, Wood Dale, IL, USA) for 5 min.

Y-maze. Spatial working memory was assessed in mice using the Y-maze (Y-shaped plastic maze, with 40 \times 20 \times 10 cm³ arm sizes). During a 5-min trial, the sequence of arm entries was recorded using the ANY-maze Video Tracking System (Stoelting). The percentage alternation was calculated as the ratio of actual to possible alternations (defined as the total number of arm entries – 2) \times 100%.

Barnes maze. The Barnes maze was used to assess hippocampus-dependent spatial learning and memory.³⁰ The Barnes maze consists of a gray, acrylic, circular disk 91 cm in diameter with 20 equally spaced holes (5 cm in diameter) located 1.5 cm from the edge, and elevated 90 cm above the floor. The platform was illuminated by a 100 W electric bulb located 110 cm above the center of the maze

(150 lx), thus providing motivation for the animals to avoid the open surface and escape into a small dark recessed chamber (escape box) located under the platform. The inclusion of false boxes that look the same as the target escape box, but are too small to be entered, were used to remove visual cues that might allow the mouse to discriminate the location of the escape hole from the other holes. A cylindrical plastic start chamber (10.5 cm diameter, 9.5 cm height) was used to hold mice in the middle of the maze at the start of each trial. A web camera (Logitech, Zurich, Switzerland) was placed 1.2 m above the center of the maze to record trials using the ANY-maze Video Tracking System (Stoelting). Distinct spatial cues placed in a constant location around the maze served as a reference point to learn the position of the escape hole.

Mice completed 4 days of acquisition training with four trials per day with an inter-trial interval of 10–15 min. For each trial, mice were placed in the start chamber in the middle of the maze, and after 10 s, the start chamber was raised to start the trial and the mouse was allowed to explore the maze for 3 min. During these 3 min, the latency to enter the escape box, distance traveled, speed, time spent immobile and search strategy used was measured, among other parameters. The trial ended when the mouse successfully entered the escape box or after the 3 min had elapsed. If a mouse did not successfully enter the escape box during the 3 min, it was placed back at the start position and gently guided to the escape hole, where the mouse remained in the escape box for 1 min. At the end of each trial, mice were returned to their holding cage until the next trial. To reduce intramaze odor cues, the maze surface and escape box were cleaned with 70% ethanol after each trial. Search strategies were determined by examining individual track plots for each mouse per day and classifying their ability to find the escape location as either: (1) random search strategy—search patterns that cross through the center of the maze in a completely random manner; (2) serial search strategy—in which mice search every hole or every other hole in a clockwise or counter-clockwise systematic manner; or (3) spatial search strategy—in which the mice were able to locate the escape box directly plus or minus two adjacent holes within the target quadrant only.

Statistical analysis. Data analysis was performed using the Statistica 10.0 (StatSoft Inc, Tulsa, OK, USA) and SPSS 19.0 (16M, Schweiz, Zurich, Switzerland) statistical software. Tests for normal distribution were performed before statistical testing, according to the results of the Shapiro–Wilk and the Kolmogorov–Smirnov test for normality, either Student's *t*-test or Mann–Whitney *U*-test for two sample groups or analysis of variance was performed (followed by *post-hoc* Fischer's least significant difference analysis). A *P*-value <0.05 was considered statistically significant. Error bars are s.e.m.

Results

Transgene expression and APP processing. The newly generated E22 Δ A β mice overexpress the human APP695 isoform containing the Swedish (K670N+M671L) and Osaka (E693 Δ) mutations. The Swedish double mutation

was introduced to increase the amount of total secreted A β without affecting the A β 42/A β 40 ratio. Several transgenic founder lines were analyzed for brain expression of the full-length human APP transgene, and a line expressing transgene levels comparable to our APP transgenic arcA β mouse model (expressing APP with the Swedish and Arctic (E693G) mutations) and the widely established Tg2576 AD mouse model (expressing APP with the Swedish mutation alone) was chosen for further analysis (Figure 1a). Western blot analysis of SDS extracts revealed similar full-length APP and β -stub (C99) levels in the E22 Δ A β , arcA β and Tg2576 mice (Figure 1a). α -Stub (C83) levels, however, were significantly reduced in both E22 Δ A β and arcA β mice as compared with Tg2576 mice (Figure 1a). MSD assay analysis of soluble APP levels from SDS brain extracts revealed a two- to threefold reduction in sAPP α levels in the E22 Δ A β and arcA β mice. In contrast, sAPP β levels were comparable to the Tg2576 mice (Figure 1b). These findings indicate that both intra-A β APP mutations at position 22 of the A β sequence specifically interfere with the α -secretase processing of APP *in vivo*.

Age-dependent changes in A β levels. As a next step, we used MSD technology to determine A β levels in four sequentially extracted protein fractions from the cortical brain tissue (Figure 2). In agreement with previous findings,^{27,31,32} Tg2576 and arcA β mice showed an age-dependent exponential accumulation of A β in detergent-insoluble (FA-soluble) protein fractions (Figure 2e–k) that accompanied the occurrence of parenchymal amyloid deposits in these mouse models (data not shown). A β accumulation occurred earlier and was more pronounced in the arcA β mice as compared with age-matched Tg2576 mice (Figure 2e–k), as previously reported in immunohistological findings.^{27,31} In contrast, most of the A β in the E22 Δ A β mice accumulated in the detergent-soluble (SDS) protein fraction up to the age of 15 months (Figure 2a–c). MSD analysis of Tris buffer and mild detergent (Triton X-100) extracts in the E22 Δ A β mice revealed only very low A β levels as compared with the Tg2576 mice (Figure 2a–c). In addition, E22 Δ A β mice (similar to the arcA β mice) showed four- to fivefold lower plasma A β levels than the Tg2576 mice (Supplementary Figure 1). These results indicate that the lack of accumulation of detergent-insoluble A β in the E22 Δ A β mice up to the age of 15 months was not accompanied by a relative increase in soluble brain or peripheral (plasma) A β levels. Western blot analysis of the four protein fractions at 15 months further confirmed the MSD findings (Supplementary Figure 2). At 24 months of age, however, a significant increase in detergent-insoluble (FA-soluble) A β was observed in the E22 Δ A β mice (Figure 2d). This change in A β solubility provided the first biochemical evidence for a substantial accumulation of fibrillar amyloid deposits in aged E22 Δ A β mice, even though FA-soluble A β levels remained relatively low in comparison to amyloid depositing Tg2576 and arcA β mice at 15 months (Figure 2b and c).

Late CAA in the absence of amyloid plaque deposition. Immunostainings of paraffin-embedded brain sections from aged E22 Δ A β mice with A β -specific antibodies showed no

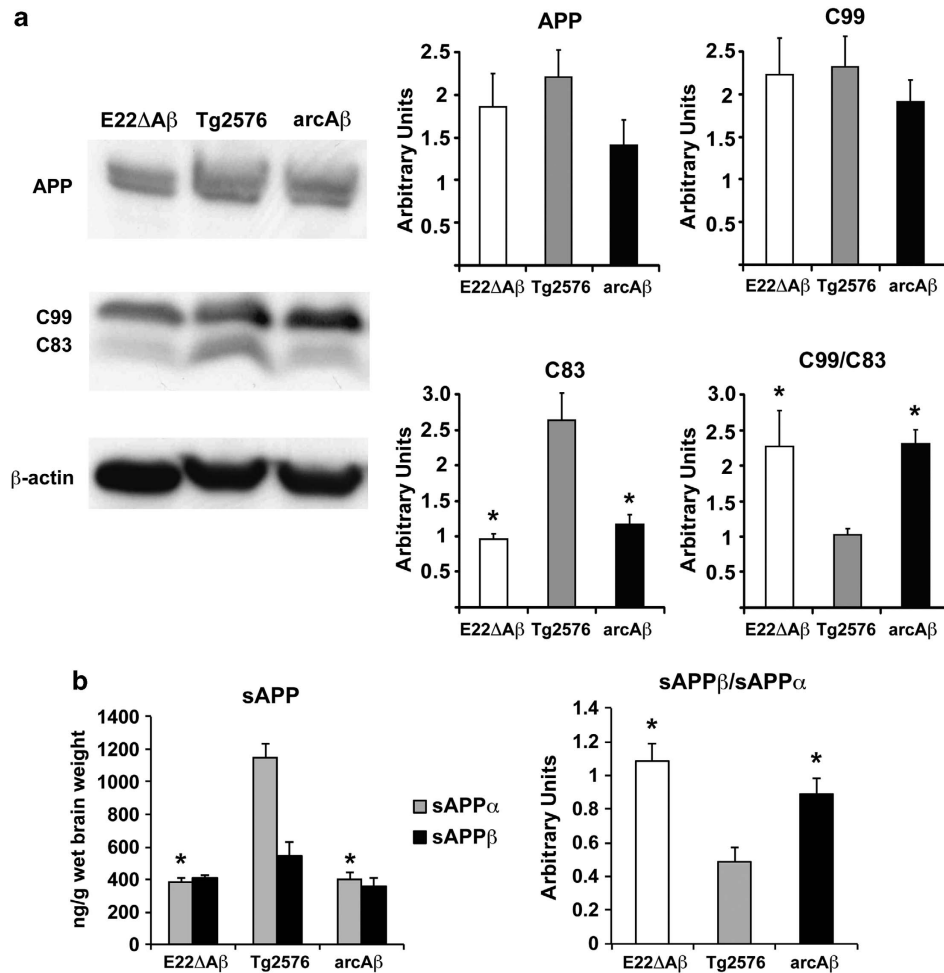


Figure 1 Intra-amyloid β (A β) mutations decrease α -cleavage *in vivo*. Transgene expression and A β peptide precursor (APP) processing were compared in three mouse models of Alzheimer's disease (AD). (a) Western blot analysis of cortical SDS extracts from 3-month-old E22ΔA β mice in comparison with age-matched Tg2576 and arcA β mice. Full-length APP and APP C-terminal fragments (C99 and C83) were detected with a C-terminal-specific APP antibody and values normalized to β -actin. Quantification reveals significant reductions in α -stub (C83) levels in both transgenic mouse lines bearing the intra-A β mutation. This results in an increased C99/C83 ratio in the E22ΔA β and arcA β mice as compared with the Tg2576 mice. * $P < 0.05$ E22ΔA β and arcA β vs Tg2576 (Mann–Whitney *U*-test). $n = 4$ per group. (b) MSD analysis of sAPP α and sAPP β levels in cortical samples extracted with Tris buffer containing 2% SDS. In accordance with the results of the APP CTF analysis, sAPP β levels were not significantly different among the three APP transgenic mouse lines, whereas sAPP α was reduced almost threefold in the E22ΔA β and arcA β mice as compared with the Tg2576 mice. * $P = 0.001$ E22ΔA β and arcA β vs Tg2576 (Mann–Whitney *U*-test). $n = 6$ –8 per group.

evidence of extracellular parenchymal amyloid deposition up to an age of 15 months (Figure 3a and b). At 24 months of age, however, extracellular vascular amyloid deposits were observed in cortical and—more pronounced—cerebellar leptomeningeal vessels (Figure 4a and b). Vascular amyloid deposits were thioflavin S (Figure 4c) and Congo red positive (Figure 4d), and were immunostained with several A β -specific antibodies (Figure 4a–f). Immunostainings with A β 40- and A β 42-specific monoclonal antibodies revealed that A β 40 was the dominant A β species deposited in the vessel walls of the E22ΔA β mice (Figure 4e and f).

Early accumulation of intracellular A β oligomers. Although extracellular (vascular) amyloid deposits were detectable in the E22ΔA β mice only as late as at the age of 24 months, immunostaining of paraffin-embedded brain

sections with the monoclonal antibody 6E10 revealed prominent intraneuronal accumulation of dot-like aggregates mainly in hippocampal CA1 (Supplementary Figure 3a) and cortical (Supplementary Figure 3b) neurons that were observed already at the age of 3 months (data not shown). Double immunostaining with a polyclonal antibody directed against the APP C terminus (Supplementary Figure 3c and d) showed little colocalization of 6E10 and APP immunoreactivity in the cortex and hippocampus (Supplementary Figure 3e and f), thus excluding 6E10-positive aggregates as full-length APP or APP C-terminal fragments. Immunostaining with A β -specific antibodies directed against the C terminus of A β 40 and A β 42 confirmed that the intraneuronal deposits indeed corresponded to A β aggregates (Supplementary Figure 4a–f). Interestingly, immunostaining with the A β 40-specific antibody led to a more diffuse staining of both dot-like aggregates and neuronal cell bodies and

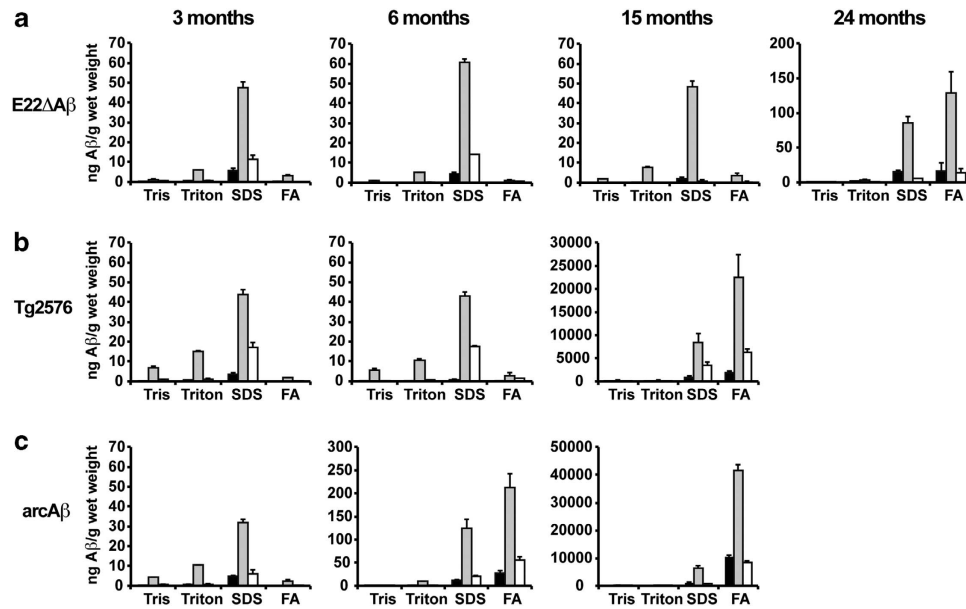


Figure 2 Accumulation of detergent-insoluble amyloid β (A β) at different ages in the three Alzheimer's disease (AD) mouse models. MesoScale Discovery (MSD) analysis of A β 38 (black column), A β 40 (gray column) and A β 42 (white column) in serially extracted cortical samples (Tris buffer, Tris buffer containing 1% Triton, Tris buffer containing 2% sodium dodecyl sulfate (SDS) and formic acid (FA)). (a) In E22 Δ A β mice most of the A β can be detected in the SDS-soluble protein fraction up to the age of 15 months, whereas Tris buffer-soluble, Triton-soluble and FA-soluble A β levels are comparatively low. Only at 24 months of age is detergent-insoluble A β detected. (b) In contrast, Tg2576 mice show higher soluble A β levels in Tris and Triton fractions and a prominent age-dependent A β accumulation in the detergent-insoluble FA fraction at 15 months. (c) The 3-month-old arcA β mice show intermediate Tris- and Triton-soluble A β levels (as compared with age-matched E22 Δ A β and Tg2576 mice) and accumulation of A β in the FA fraction already at the age of 6 months. $n = 6-9$ per group.

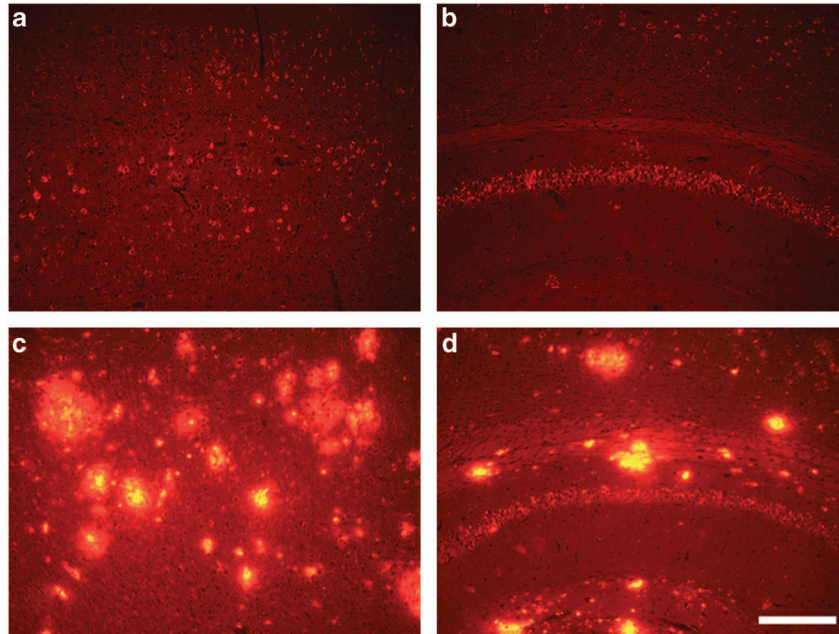


Figure 3 Lack of amyloid deposition in 15-month-old E22 Δ A β mice. (a–d) Immunostaining with antibody 6E10 reveals a complete absence of extracellular amyloid deposits in 15-month-old E22 Δ A β mice (a: cortex; b: hippocampus). In contrast, arcA β mice show a massive extracellular amyloid deposition at that age (c: cortex; d: hippocampus). Scale bars: 250 μ m (a–d). A β , amyloid β .

processes (Supplementary Figure 4a, b and e), whereas the A β 42-specific specifically stained compact intraneuronal aggregates (Supplementary Figure 4c, d and f).

Further characterization of the intraneuronal A β aggregates in the E22 Δ A β mice with several oligomer-specific (conformation-dependent) antibodies revealed that the dot-like A β

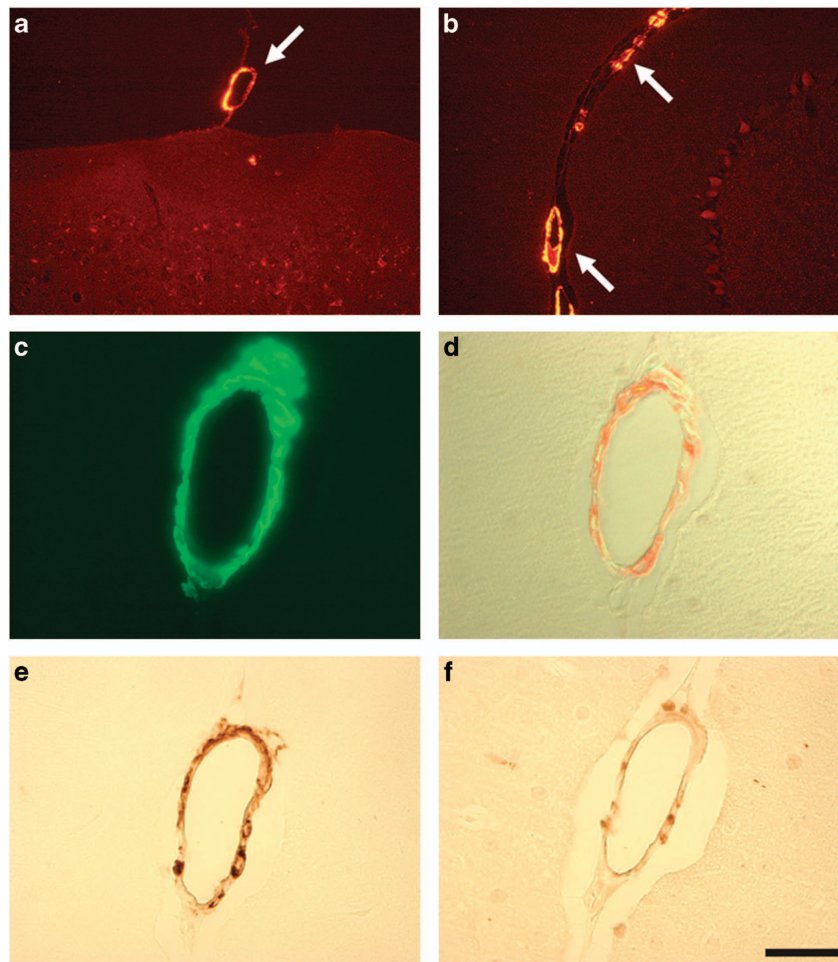


Figure 4 Aged E22 Δ A β mice develop congophilic amyloid angiopathy (CAA) at 24 months. (a + b) Immunostaining of amyloid-laden leptomeningeal vessels with polyclonal anti-amyloid antibody directed against human pan-A β in the cortex (a) and cerebellum (b) of a 24-month-old E22 Δ A β mouse. (c + d) Vascular amyloid deposits are thioflavin S (c) and Congo red positive (d). Note the classical apple-green birefringence of Congo red-stained vessels under polarized light (d). (e + f) Immunostainings with A β 40-specific monoclonal antibody BA27 (e) and A β 42-specific monoclonal antibody BC05 (f) reveal that A β 40 is the dominant A β species deposited in the vessel walls. Scale bars: 125 μ m (a + b) and 40 μ m (c–f). A β , amyloid β .

deposits were strongly stained by the polyclonal OC antibody directed against fibrillar A β oligomers³³ (Figure 5a–d), but not the A11 antibody directed against prefibrillar oligomers³⁴ (Supplementary Figure 4g and h). OC-immunoreactive intraneuronal deposits were observed in CA1 hippocampal and cortical neurons as early as 3 months of age and appeared to grow and become more compact as the mice aged (Figure 5a–d). Strikingly, the intraneuronal oligomeric A β deposits were also recognized by 11A1 (Figure 5e and f), a novel conformation-dependent monoclonal antibody specifically designed against the turn between positions 22 and 23 of the A β sequence. 11A1 has recently been shown to stain intraneuronal A β aggregates in brains of AD patients but not APP transgenic mice.²⁶

Early cognitive deficits in transgenic E22 Δ A β mice. Based on our previous findings in the arcA β mouse model in which intracellular A β deposits occurred concomitantly with robust cognitive deficits, we hypothesized that the early accumulation of intraneuronal oligomeric A β deposits,

particularly in hippocampal brain regions of the E22 Δ A β mice, would also be accompanied by significant impairments on several cognitive tasks. Both E22 Δ A β mice and wild-type littermates displayed, across all age groups, similar general health measures, auditory–visual sensory integrity, comparable grip strength, intact righting and extension reflexes. Assessment of body weight demonstrated equivalent body weights across all experimental groups, for both males and females. Locomotor and anxiety examination using the open field test demonstrated the published phenotype of increased locomotor activity and exploratory behavior, including increased total distance traveled, less time immobile and more time spent in the center zone, in transgenic mice (data not shown). Cognitive assessment in the Y-maze demonstrated a significant decrease in the percentage spatial alternation rate for E22 Δ A β mice at 3 months of age ($t_{(55)} = -3.16$, $P < 0.005$), 6 months of age ($t_{(28)} = -1.727$, $P < 0.05$) and 9 months of age ($t_{(16)} = -1.772$, $P < 0.05$), as compared with their wild-type littermates (Figure 6a). The Barnes maze was also used at 3 months of age as a second

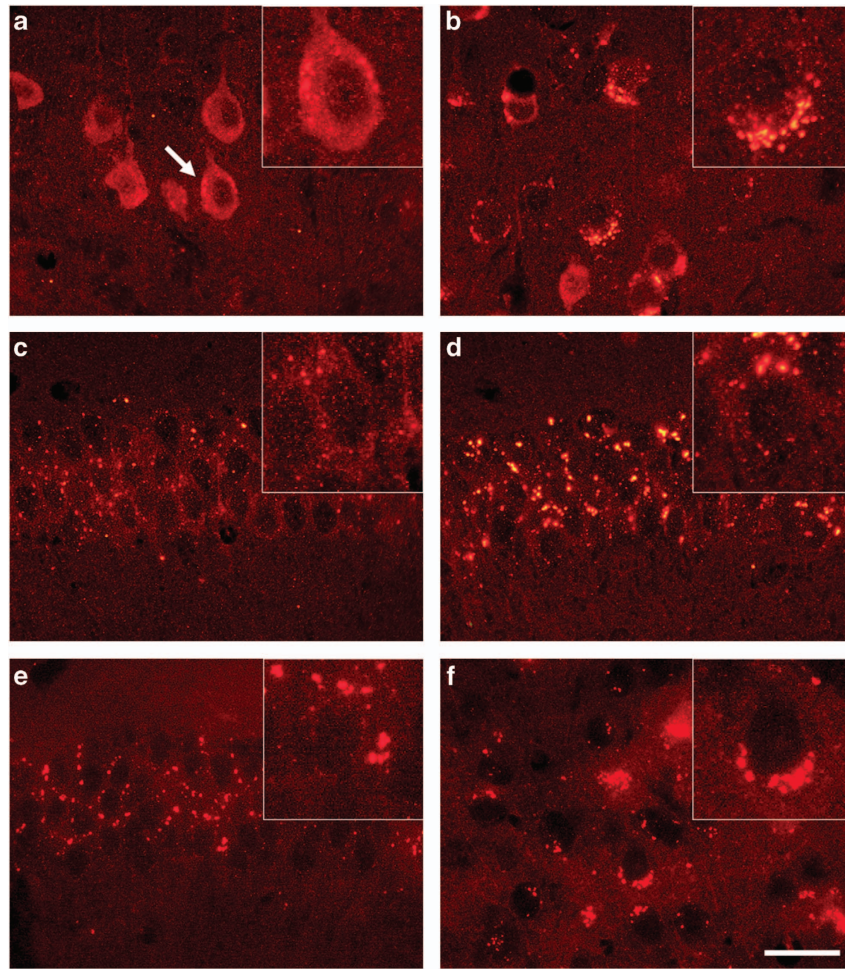


Figure 5 Intraneuronal amyloid β (A β) consists of fibrillar oligomers bearing a 'toxic turn' conformation. (a–d) Immunostaining with polyclonal OC antibody directed against fibrillar A β oligomers reveals age-dependent accumulation of intracellular oligomeric deposits in cortical (a and b) and CA1 hippocampal neurons (c and d). Note the temporal change of intraneuronal oligomers from rather diffuse and smaller dot-like aggregates at 3 months of age (a and c) to bigger and more compact deposits at 15 months of age (b and d). (e and f) Conformation-dependent monoclonal antibody 11A1 directed against the turn between positions 22 and 23 of the A β sequence stains hippocampal (e) and cortical (f) intraneuronal aggregates (representative immunostaining in a 15-month-old E22 Δ A β mouse). Scale bar: 40 μ m (a–f).

hippocampus-dependent cognitive task to assess spatial learning and memory. Results demonstrated that both wild-type and transgenic E22 Δ A β mice showed successful learning, with significantly lower latencies to escape over the 4 days of training (all P 's < 0.05; Figure 6b). Between genotypes, a significant main effect was shown for E22 Δ A β mice ($F_{(1,340)} = 16.656$, $P < 0.001$). Fischer's least significant difference *post hoc* analysis demonstrated a significantly longer latency to escape for E22 Δ A β mice, as compared with wild-type littermates on day 3 ($P = 0.01$) and day 4 ($P = 0.03$), with similar but nonsignificant trends on day 1 ($P = 0.096$) and day 2 ($P = 0.095$) (Figure 6b). Qualitative assessment of Barnes maze search strategies revealed a decrease in spatial search strategies in 3-month-old E22 Δ A β mice (Figure 6d), as compared with wild-type control mice (Figure 6c).

Unique aggregation properties of recombinant E22 Δ -mutated A β peptides. In agreement with our previous findings that both E22 Δ A β 40 and E22 Δ A β 42 readily formed amyloid fibrils *in vitro*, we observed fibrillar (congophilic)

amyloid deposits *in vivo* in aged APP transgenic E22 Δ A β mice. Extracellular amyloid deposition in the E22 Δ A β mice, however, only occurred as leptomeningeal CAA at an advanced age of 24 months (Figure 4), whereas wild-type A β -producing Tg2576 mice and E22G (Arctic) A β -producing arcA β mice accumulated fibrillar amyloid deposits at a much earlier age (Figures 2 and 3).^{27,31} To further investigate the biophysical basis of the E22 Δ intra-A β mutation leading to early intracellular and very late extracellular amyloid deposition *in vivo*, we conducted *in vitro* thioflavin T aggregation assays of recombinant E22 Δ A β 40 and E22 Δ A β 42 peptides. E22 Δ A β aggregation curves were compared with the aggregation curves of A β 40 and A β 42 with wild-type or Arctic sequence (Figure 7). Measurements were terminated as the thioflavin T signal reached a plateau, after which the different A β preparations were further incubated for a total of 1 h, followed by western blot analysis. Results demonstrated typical aggregation kinetics of wild-type A β 42 and E22G A β 42 peptides involving a lag phase, an exponential growth phase and a plateau phase of saturated fibril growth

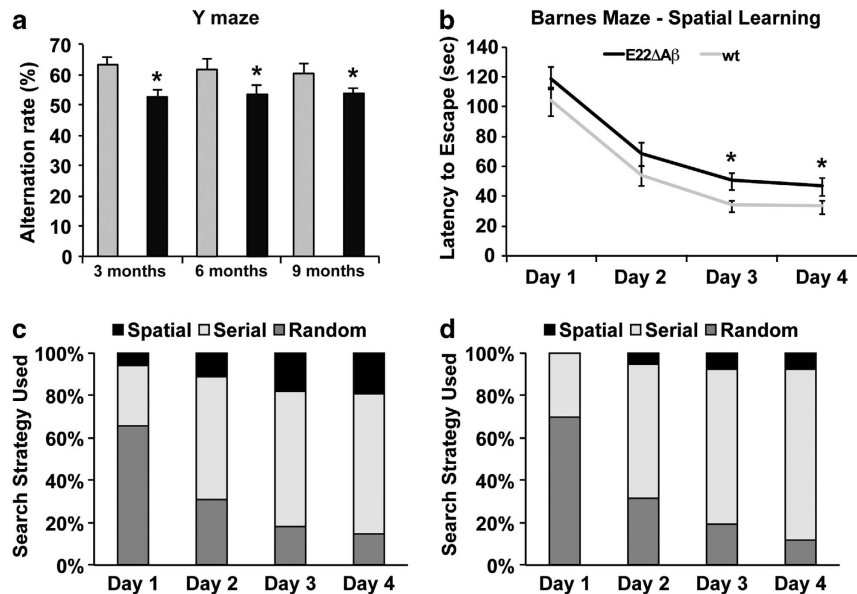


Figure 6 Early cognitive deficits in E22ΔA β mice. (a) Cognitive assessment in the Y-maze demonstrates a significant decrease in the percentage spatial alternation rate for E22ΔA β mice as compared with their wild-type (wt) littermates at 3, 6 and 9 months of age ($P < 0.05$ for all comparisons; Student's *t*-test). (b) Assessment of Barnes maze spatial learning at the age of 3 months reveals that both wild-type and E22ΔA β transgenic mice show successful learning over the 4 days of training. However, E22ΔA β mice, as compared with wild-type littermates, show significantly higher latencies to escape on day 3 ($P = 0.01$) and day 4 ($P = 0.03$), with similar but nonsignificant trends on day 1 ($P = 0.096$) and day 2 ($P = 0.095$) (analysis of variance (ANOVA), followed by *post hoc* Fisher's least significant difference (LSD) analysis). (c and d) Qualitative assessment of Barnes maze search strategies reveals a decrease in spatial and relative increase in serial search strategies in 3-month-old E22ΔA β mice (d) in comparison to wild-type control mice (c). $n = 22$ – 24 per group. A β , amyloid β .

(Figure 7a and b). In contrast, no obvious lag phase and no exponential growth phase were observed for E22Δ A β 42 at $2.5 \mu\text{M}$ (Figure 7a) and—more obvious—at $5 \mu\text{M}$ initial concentration (Figure 7b). Instead, E22Δ A β 42 showed a constant non-exponential increase in thioflavin T fluorescence from time point 0 on and—in contrast to E22G and wild-type A β 42—formed detergent-insoluble high-molecular-weight aggregates immediately after reconstitution at pH 7.4, as revealed by western blotting of the recombinant peptide preparations that were used for the thioflavin T analysis (Figure 7c). Thioflavin T analysis of recombinant A β 40 peptide variants revealed a dramatically accelerated fibril formation for both E22Δ and E22G A β 40 in comparison to wild-type A β 40 (Figure 7d). In contrast to E22Δ A β 42, E22Δ A β 40 aggregation was characterized by a lag phase, an exponential growth phase and a plateau phase with unusually high absolute thioflavin T fluorescence values indicating increased thioflavin T binding capacity of the E22Δ A β 40 peptide as reported previously.²⁴

In summary, thioflavin T analysis of recombinant E22Δ A β 42 and E22Δ A β 40 aggregation kinetics confirmed our previous findings of a highly increased fibrillogenic property of single preparations of the two peptide variants.²⁴

Inhibition of E22ΔA β aggregation in the presence of wild-type A β . The E693Δ APP mutation is one of the two currently known familial AD mutations in the *APP* gene with a recessive Mendelian trait of inheritance.^{9,35} The second recessive mutation, A673V, has recently been shown to be highly amyloidogenic in the homozygous state, but anti-amyloidogenic in the heterozygous state as revealed by an

inhibition of A β aggregation when mutated and wild-type peptides were co-incubated.³⁵ We hypothesized similar effects on the aggregation kinetics when incubating mixtures of the wild-type and E22Δ-mutated A β peptides. Co-aggregation of wild-type A β 42 with E22Δ A β 42 led to a significant extension of the lag phase and a delay of the growth phase in comparison to the aggregation of wild-type A β 42 alone. This effect was specific for the E22Δ intra-A β mutation, as it was not observed with the E22G A β 42 variant. In contrast, co-aggregation of E22G (Arctic) A β 42 with wild-type A β 42 revealed a similar aggregation kinetics as E22G A β 42 alone (see Figure 8a). Similarly, the aggregation of E22Δ A β 40 peptides was also significantly delayed in mixtures containing 50% wild-type A β 40, whereas co-incubation of E22G A β 40 with wild-type A β 40 only slightly delayed fibril formation in comparison to E22G A β 40 alone (Figure 8b). In conclusion, only the E22Δ intra-A β mutation (but not the E22G variant) was associated with a significant delay of aggregation in the heterozygous state, simulated by equal mixtures with wild-type A β peptides.

Inhibition of E22Δ A β 40 aggregation in the presence of E22Δ A β 42. Based on our results that co-incubation of wild-type A β with E22ΔA β significantly delayed amyloid fibril formation, we hypothesized that co-incubation of E22Δ A β 40 with E22Δ A β 42 may have similar antifibrillogenic effects. Therefore, recombinant E22Δ A β 40 and E22Δ A β 42 peptides were co-incubated in a physiological 9:1 ratio at $2.5 \mu\text{M}$, and the increase in thioflavin T fluorescence was monitored over time (Figure 8c). We found a dramatic delay of the growth phase and a remarkable loss of absolute

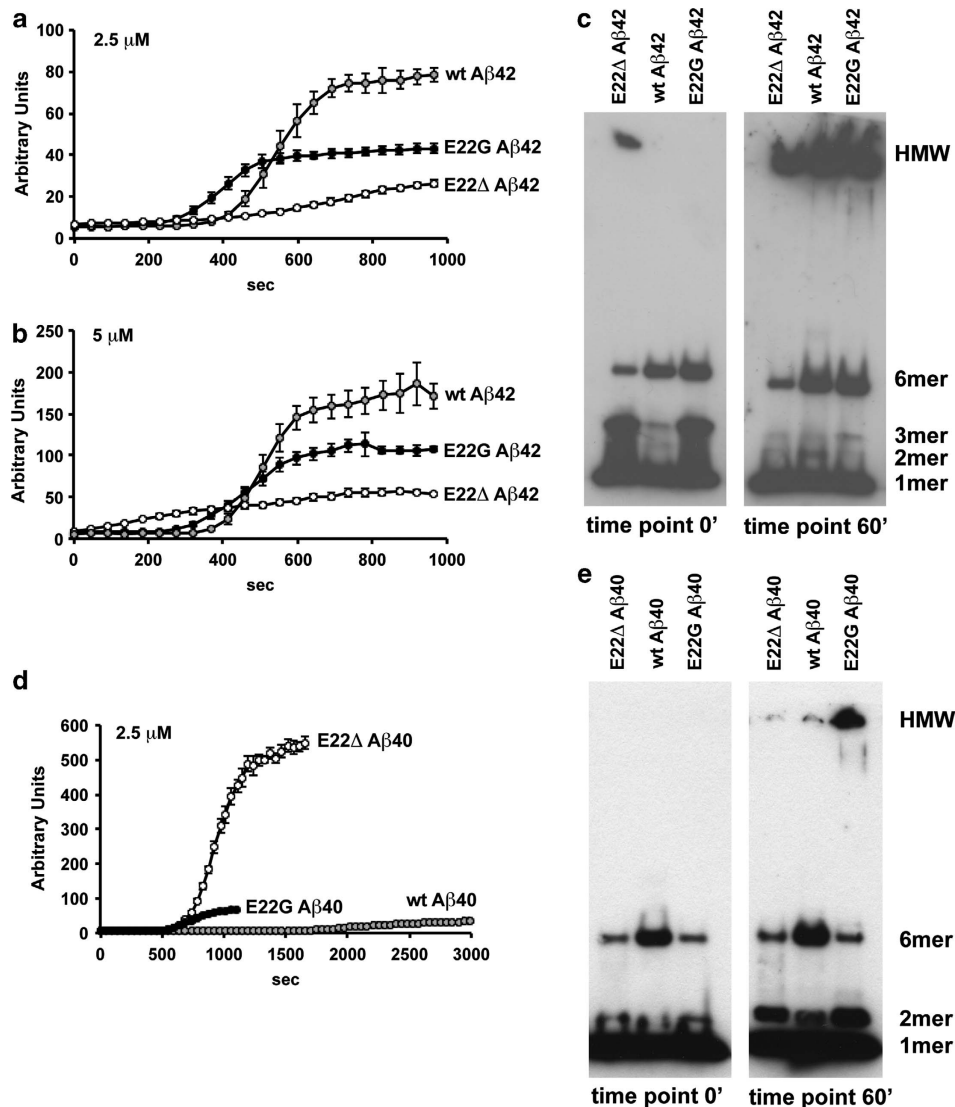


Figure 7 Unique aggregation kinetics of recombinant E22 Δ A β peptides *in vitro*. Fibril formation is measured via the absolute increase in thioflavin T fluorescence at 482 nm during aggregation. Aggregation of amyloid β (A β) peptides (identical total monomer concentrations: 2.5 μ M in **a** + **c**, 5 μ M in **b**) at pH 7.4 and 37 °C is initiated by a 10-fold dilution of A β stock solutions in 10 mM NaOH with buffer. Average curves of three independent experiments are shown (\pm s.e.m.). (**a** and **b**) Thioflavin T analysis of E22G (Arctic) A β 42 and wild-type (wt) A β 42 reveals that a typical aggregation kinetics with a lag phase, which is shorter for E22G A β 42 than wild-type A β 42, is followed by an exponential growth phase that eventually reaches a plateau of saturated β -sheet formation. In contrast, no obvious lag phase and no exponential growth phase is observed for E22 Δ A β 42 at 2.5 μ M (**a**) and—more obvious—at 5 μ M (**b**). Instead, E22 Δ A β 42 is characterized by slow non-exponential fibril growth; however, it forms detergent-insoluble high molecular weight (HMW) aggregates immediately after reconstitution at pH 7.4 as revealed by western blotting of the recombinant peptide preparations that were used for the thioflavin T analysis at time point 0 and time point 60 of the assay (**c**). (**d**) In contrast to E22 Δ A β 42, E22 Δ A β 40—similar to E22G A β 40—shows an aggregation kinetics characterized by a lag phase, an exponential growth phase and a plateau phase. Note the unusually high thioflavin T fluorescence values for E22 Δ A β 40 in contrast to E22G and wild-type A β 40, and the remarkably shorter lag phases of the E22-mutated A β 40 variants in comparison to the wild-type peptide. (**e**) In contrast to the amyloid β (A β 42) analysis, western blot analysis of the A β 40 peptide variants reveals no immediate formation of detergent-insoluble HMW aggregates at time point 0 min. Only the E22G mutant develops prominent HMW aggregates after 60 min.

thioflavin T fluorescence when we compared the aggregation curve with the aggregation curve of E22 Δ A β 40 alone (Figures 7d and 8c). Again in contrast, 9:1 mixtures of E22G A β 40 and E22G A β 42 resulted in rapid aggregation, similar to E22G A β 40 alone (Figures 7c and 8c). These results again reveal the strong aggregating properties of the E22G mutation and the inhibitory effect of the E22 Δ mutation under conditions when different peptide species coexist, as is the case *in vivo*.

Poor seeding of wild-type A β 42 aggregation by E22 Δ A β 42 fibrils. Based on the assumption that amyloid formation can be seeded by preformed amyloid fibrils, we finally addressed the question whether preaggregated E22 Δ A β 42 fibrils were capable of (a) seeding their own growth and (b) seeding the growth of wild-type A β 42 fibrils. As control experiments, wild-type A β 42 and E22G (Arctic) A β 42 fibrils were incubated with preparations of their respective monomeric peptides, which resulted in an immediate increase in

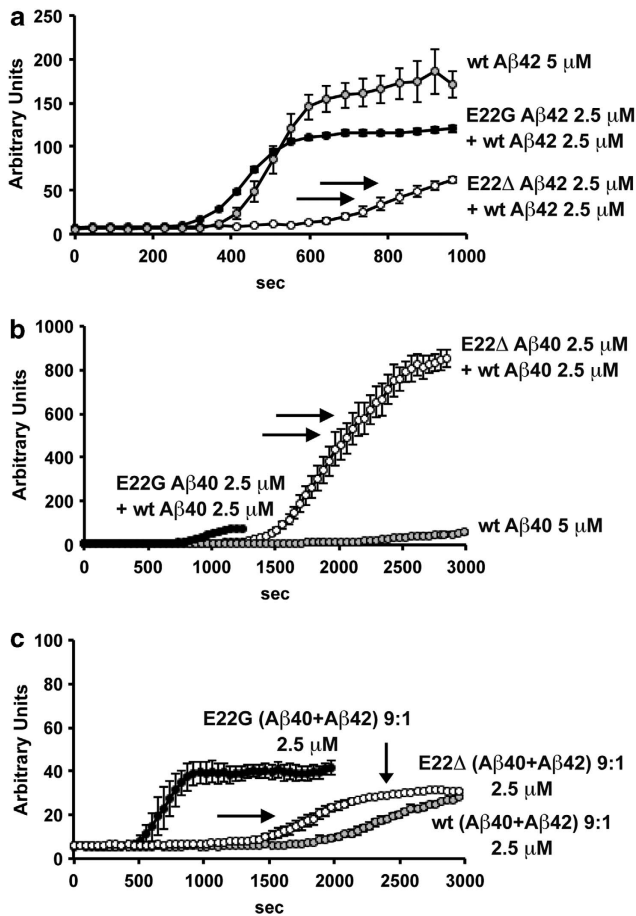


Figure 8 Anti-amyloidogenic property of E22 Δ A β peptides in mixtures of different A β peptide variants. (a and b) Co-aggregation of E22 Δ A β 42 with wild-type (wt) A β 42 (a) or E22 Δ A β 40 with wild-type A β 40 (b) leads to a significant extension of the lag phase and a delay of the growth phase in comparison with the aggregation of the respective A β peptide variants alone (see Figure 7) (total monomer concentrations for each peptide variant: 2.5 μ M). In contrast, co-aggregation of 2.5 μ M E22G (Arctic) A β 42 with 2.5 μ M wild-type A β 42 (a) results in a similar aggregation curve as with 2.5 or 5 μ M E22G A β 42 alone (see Figure 7). Co-aggregation of E22G A β 40 with wild-type A β 40 (b) only slightly delays β -sheet formation in comparison with E22G A β 40 alone. (c) Co-aggregation of E22 Δ A β 40 with E22 Δ A β 42 in a physiological 9:1 ratio (total monomer end concentration in solution: 2.5 μ M) leads to a strong inhibition of E22 Δ A β 40 aggregation. Note the dramatic delay of the growth phase and remarkable loss of absolute thioflavin T fluorescence in comparison to E22 Δ A β 40 alone (Figure 7). In contrast, 9:1 mixtures of E22G A β 40 and E22G A β 42 result in an aggregation curve very similar to E22G A β 40 alone. Average curves of three independent experiments are shown (\pm s.e.m.).

thioflavin T fluorescence without a lag phase, thus indicating good seeding properties (Supplementary Figure 4a and b). Incubation of preaggregated E22 Δ A β 42 fibrils with fresh preparations of E22 Δ A β 42 peptide also resulted in an immediate increase in thioflavin T fluorescence that was steeper as compared with the aggregation curve of E22 Δ A β 42 in the absence of fibrillar seeds (Figure 7a and Supplementary Figure 5c). However, when E22 Δ A β 42 fibrils were used for the seeding of wild-type A β 42, a significant delay of fibril formation was observed, which indicated poor seeding properties (Supplementary Figure 5e). In contrast,

E22G A β 42 fibrils appeared only slightly inferior to wild-type A β 42 fibrils in seeding wild-type A β 42 aggregation (Supplementary Figure 5d).

Discussion

Homozygous E693 Δ APP mutation carriers develop an AD-like clinical phenotype characterized by early memory disturbances, visuospatial deficits and executive dysfunction, followed by atypical neurological signs including cerebellar ataxia and gait difficulties during later stages of the disease.^{9,36} Brain amyloid imaging by Pittsburgh Compound B positron emission tomography revealed a very weak signal in E693 Δ APP mutation carriers with advanced dementia, suggesting that AD-like clinical symptoms occurred in the absence of relevant amyloid deposition in these patients.^{9,36} *In vitro* experiments with synthetic E22 Δ A β 40 and E22 Δ A β 42 peptide preparations demonstrated an enhanced oligomerization propensity, but no fibril formation,⁹ and led to the hypothesis that AD-like dementia in patients can be caused by the sole presence of synaptotoxic A β oligomers, which was well in line with previous experimental findings.^{37–43} In contrast to these initial *in vitro* findings, subsequent work, including our own recent study with highly pure recombinant peptide preparations, demonstrated that both E22 Δ A β 40 and E22 Δ A β 42 readily formed amyloid fibrils *in vitro*.^{20,24,44} Based on these *in vitro* findings, we hypothesized that E22 Δ A β 40 and E22 Δ A β 42 would, at least in principle, also form amyloid fibrils *in vivo*. Indeed, aged E22 Δ A β mice accumulated detergent-insoluble A β and showed thioflavin S and Congo red-positive amyloid deposits in leptomeningeal cortical and—more pronounced—cerebellar vessels; in contrast to the recently published E693 Δ mouse model, which completely lacked extracellular amyloid deposition even at an age of 24 months,²² E22 Δ A β mice overexpress human APP at levels comparable to those in the Tg2576 mouse line, and the introduction of the Swedish double mutation results in an additional increase in total A β levels.⁶ Taken together, this might explain the differences between our model and the E693 Δ mice. Interestingly, CAA deposits in the E22 Δ A β mice were more pronounced in the leptomeningeal vasculature of the cerebellum than in cortical vessels, which is in agreement with recent amyloid positron emission tomography findings in E693 Δ mutation carriers showing a relative increase in Pittsburgh Compound B retention in cerebellar brain regions.³⁶ The identification of CAA as a key neuropathological feature of aged E22 Δ A β mice adds the Osaka E693 Δ mutation to the list of vasculotropic intra-A β APP mutations essentially comprising all of the currently known mutations at or around position 22 of the A β sequence (for a review see Kumar-Singh¹⁸). Our immunohistological analysis of CAA deposits identified A β 40 as the major E22 Δ A β species depositing in the vessel walls of the E22 Δ A β mice, which is in line with previous findings in sporadic and familial AD cases,^{45,46} including the Dutch E693Q APP mutation.⁴⁷ Further experiments are needed to elucidate the mechanisms underlying the vasculotropism of E22 Δ -mutated A β and whether similar mechanisms as previously reported for the Dutch mutation (that is, a reduced receptor-mediated clearance across the blood–brain barrier) have a role.^{48,49}

E22 Δ A β mice—similar to our previously reported arcA β mice²⁷—develop intraneuronal A β aggregates that coincide with cognitive deficits beginning at 3 months of age. Intraneuronal A β accumulation is generally believed to be an early event in AD pathogenesis, although its relevance and role in the disease process remain a controversial topic.^{50–52} This may be partly due to technical considerations, including the use of nonspecific antibodies for the detection of intraneuronal A β . The intraneuronal A β staining in our E22 Δ A β mouse model only partially colocalized with staining of the APP C terminus, excluding false interpretation of A β staining emerging from full-length APP or APP C-terminal fragments. Furthermore, intraneuronal A β was stained by different A β C-terminus-specific antibodies, which do not crossreact with APP or APP fragments, including BC05 and BA27.⁵³ These results therefore indicate that the intraneuronal aggregates in the E22 Δ A β mice indeed correspond to A β and not accumulating APP/APP fragments. The assembly state of the intraneuronal aggregates was characterized by immunohistochemistry using several oligomer-specific antibodies. Accumulation of intracellular oligomeric A β species has recently been reported in other APP transgenic mouse lines, including the McGill-Thy1-APP mice,⁵⁴ the 3 \times Tg-AD⁵⁵ and the E693 Δ mice.²² Further immunohistological characterization of the intraneuronal deposits in the E22 Δ A β mice revealed that the intraneuronal aggregates were strongly stained by OC antibody, directed against fibrillar oligomers, but not A11 antibody, which recognizes prefibrillar oligomers.^{33,34} Interestingly, soluble fibrillar oligomers detected by OC antibody (but not prefibrillar oligomers) have recently been shown to be elevated in multiple brain regions of AD patients and to correlate with cognitive dysfunction.⁵⁶ Building on the hypothesis that intra-A β mutations at or around position 22 of the A β increase A β fibrillogenesis through a facilitation of a ‘toxic turn’ conformation between positions 22 and 23 of the A β sequence,⁵⁷ Murakami *et al.*²⁶ developed a novel monoclonal antibody (11A1) directed against this specific conformational epitope of A β 42. In human AD brains, 11A1 stained both extracellular and intracellular A β deposits, whereas in APP transgenic Tg2576 mice, only extracellular deposits were labeled.²⁶ Interestingly, 11A1 stained intraneuronal A β aggregates in the brains of the E22 Δ A β mice, suggesting that the oligomeric E22 Δ -mutated A β deposits contained the ‘toxic turn’ conformation.

Western blot and MSD analysis of APP cleavage products from cortical brain extracts revealed a relative increase in amyloidogenic (β -secretase-mediated) APP processing in the two mouse models bearing the E22 intra-A β APP mutations, which is in accordance with previous *in vitro* findings.^{21,58,59} As both E22 Δ A β and arcA β mice overexpress similar full-length APP, sAPP β and C99 levels, but significantly reduced sAPP α and C83 levels, as compared with the Tg2576 mice, we concluded that the two intra-A β mutations specifically interfered with the α -secretase cleavage of APP *in vivo*. Previous *in vitro* reports in the Arctic mutation revealed that E693G APP was not a poor substrate to α -secretase, but instead reduced APP levels at the cell surface making Arctic APP less available for α -secretase cleavage, and increasing A β levels, especially at intracellular locations.⁵⁸ Similar to the Arctic APP mutation, E693 Δ APP overexpression in cell

culture was also associated with reduced extracellular A β levels *in vitro*.^{21,58,59} Although these results may imply similar effects of the two intra-A β mutations on APP processing, we currently cannot exclude E693 Δ APP as an inferior substrate to α -secretase-mediated cleavage, thus leading to the relative increase in amyloidogenic APP processing, as shown previously for other intra-A β APP mutations, including the Flemish E692G APP mutation.⁶⁰

MSD analysis revealed a marked reduction of buffer- and Triton buffer-soluble brain A β , as well as peripheral plasma A β levels, in particular when E22 Δ A β mice were compared with age-matched Tg2576 mice-expressing wild-type A β . This indicated that the absence of extracellular amyloid deposition up to an age of 15 months was not associated with a relative increase in soluble brain or plasma A β levels. Instead, most of the A β accumulated in the SDS-soluble protein fraction likely corresponding to intraneuronal A β pools. These results are in agreement with recent *in vitro* results from cell culture experiments showing increased intracellular A β accumulation, but markedly reduced secreted A β levels in E693 Δ APP-transfected cell lines.^{9,21,23} The early accumulation of intracellular fibrillar oligomeric A β deposits in brains of the E22 Δ A β mice is in line with the unique aggregation profiles of the recombinant E22 Δ A β 40 and E22 Δ A β 42 peptides showing accelerated β -sheet formation in thioflavin T aggregation assays as compared with the respective wild-type A β peptides^{20,24,44} (and this study). When aggregation curves of E22 Δ , E22G and wild-type A β 40 and A β 42 were compared, E22 Δ A β 42 showed the highest fibrillogenesis propensity as it aggregated without a lag phase and formed SDS-resistant amyloid fibrils immediately after reconstitution in solution at pH 7.4. However, E22 Δ A β 42 fibril growth after reconstitution in solution occurred only in a slow, non-exponential manner in contrast to the other peptides, including E22 Δ A β 40, whose aggregation was characterized by an exponential growth phase following lag phase. This slow fibril growth and lack of exponential growth in E22 Δ A β 42 may be a crucial factor in the prevention of extracellular amyloid deposition in the E22 Δ A β mice.

Apart from A673V, a recently described familial AD APP mutation in an Italian pedigree, E693 Δ , is the second currently known recessive APP mutation that is pathogenic only in the homozygous state.^{9,35} Similar to the A673V mutation, which is anti-amyloidogenic in the heterozygous state,³⁵ our co-aggregation experiments with recombinant E22 Δ A β peptide variants also revealed an inhibition of A β aggregation when mutated and wild-type peptides were co-incubated. In contrast, co-aggregation of E22G (Arctic) A β peptides with the respective wild-type A β peptides resulted in aggregation curves very similar to those of the E22G A β peptides alone. Moreover, E22 Δ A β 42 fibrils, in contrast to wild-type A β 42 and E22G A β 42 fibrils, only very inefficiently seeded wild-type A β 42 fibrillogenesis. These results provide a possible explanation why heterozygous carriers of the E693 Δ mutation do not develop the disease, whereas heterozygous carriers of the autosomal dominant Arctic mutation develop dementia.

Hence, we propose the following hypothetical model to explain the phenotype of the Osaka E693 Δ mutation, in which E22 Δ A β aggregation occurs primarily in intracellular compartments where peptide concentrations are high enough to allow

for aggregate formation (Supplementary Figure 6). Outside the cell, E22 Δ A β peptide variants may interact with each other—and possibly also other peptides—in a way that results in an inhibition of aggregation and amyloid seed formation. Moreover, the E22 Δ A β 42 peptide shows specific aggregation properties (slow non-exponential growth, lack of exponential growth phase) that likely further prevent parenchymal amyloid plaque deposition (Supplementary Figure 6). However, late vascular amyloid deposition occurs possibly due to the vasculotropism of the E22-mutated A β peptide variant and age-related changes along perivascular clearing pathways (Supplementary Figure 6).

Conflict of interest

Noriaki Kinoshita is employee of Immuno-Biological Laboratories, Gunma, Japan. Luka Kulic, Jordan McAfoose, Tobias Welt, Christian Tackenberg, Claudia Späni, Fabian Wirth, Verena FINDER, Uwe Konietzko, Maria Giese, Anne Eckert, Takahiro Shimizu, Kazuma Murakami, Kazuhiro Irie, Suhail Rasool, Charles Glabe, Christoph Hock and Roger M Nitsch have no competing interests to declare.

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